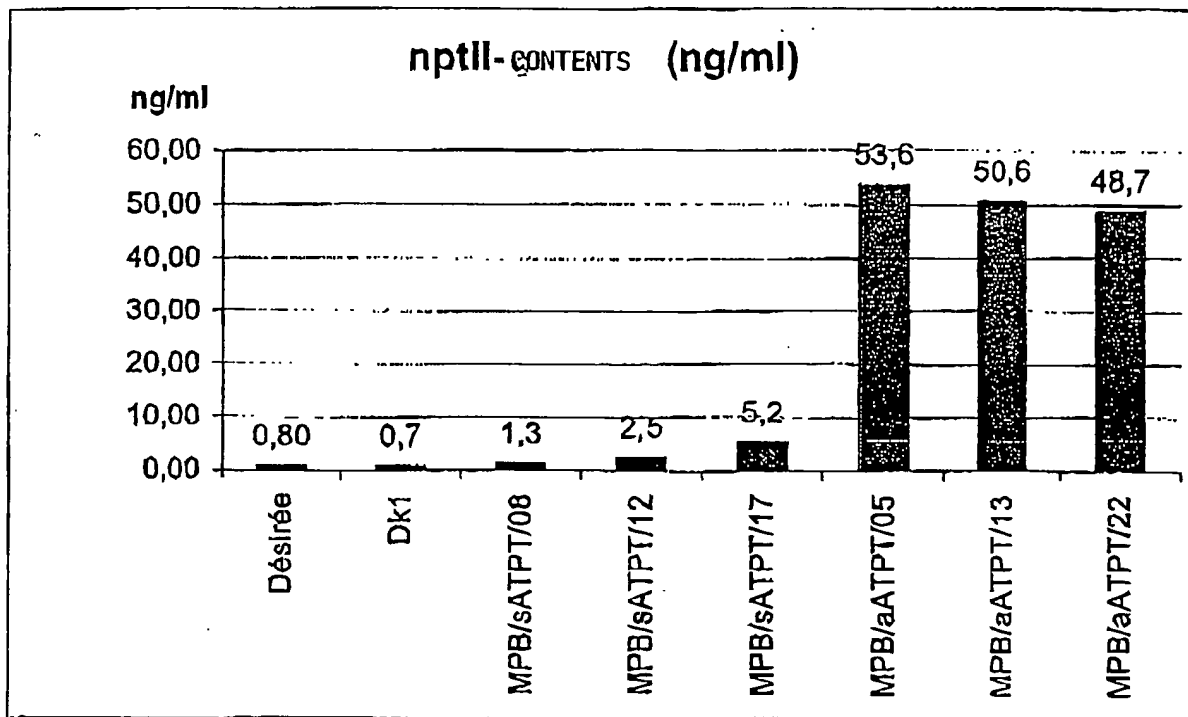


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Figur 1:



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Fig. 2

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atggtttcca aaccttttct atctttgctt tcacttttct tgccttcttt tacaagcaca	60
tgttta gca gct gat gtg cag ctg gtg gag tct ggg gga ggc tta gtg Ala Ala Asp Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val	108
1 5 10	
cag cct gga ggg tcc cgg aaa ctc tcc tgt gca gcc tct gga ttc act Gln Pro Gly Gly Ser Arg Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr	156
15 20 25 30	
ttc agt agc ttt gga atg cac tgg gtt cgt cag gct cca gag aag ggg Phe Ser Ser Phe Gly Met His Trp Val Arg Gln Ala Pro Glu Lys Gly	204
35 40 45	
ctg gag tgg gtc gca tat att agt agt ggc agt agt acc atc tac tat Leu Glu Trp Val Ala Tyr Ile Ser Ser Gly Ser Ser Thr Ile Tyr Tyr	252
50 55 60	
gca gac aca gtg aag ggc cga ttc acc atc tcc aga gac aat ccc aag Ala Asp Thr Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Pro Lys	300
65 70 75	
aac acc ctg ttc ctg caa atg acc agt cta agg tct gag gac acg gcc Asn Thr Leu Phe Leu Gln Met Thr Ser Leu Arg Ser Glu Asp Thr Ala	348
80 85 90	
atg tat tac tgc gca aga gat tac ggg gct tat tgg ggc caa ggg acc Met Tyr Tyr Cys Ala Arg Asp Tyr Gly Ala Tyr Trp Gly Gln Gly Thr	396
95 100 105 110	
acg gtc acc gtc tcc tca ggt gga ggc ggt tca ggc gga ggt ggc tct Thr Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser	444
115 120 125	
ggc ggt ggc gga tgc gac att gag ctc acc cag tct cca gca atc atg Gly Gly Gly Gly Ser Asp Ile Glu Leu Thr Gln Ser Pro Ala Ile Met	492
130 135 140	
tct gca tct cca ggg gag aag gtc acc atg acc tgc agt gcc agt tca Ser Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys Ser Ala Ser Ser	540
145 150 155	
agt gta agg tac atg aac tgg ttc caa cag aag tca ggc acc tcc ccc Ser Val Arg Tyr Met Asn Trp Phe Gln Gln Lys Ser Gly Thr Ser Pro	588
160 165 170	
aaa aga tgg att tat gac aca tcc aaa ctg tct tct gga gtc cct gct Lys Arg Trp Ile Tyr Asp Thr Ser Lys Leu Ser Ser Gly Val Pro Ala	636
175 180 185 190	
cgc ttc agt ggc agt ggg tct ggg acc tct tac tct ctc aca atc agc Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Tyr Ser Leu Thr Ile Ser	684
195 200 205	
agc atg gag gct gaa gat gct gcc act tat tac tgc cag cag tgg agt Ser Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp Ser	732
210 215 220	
agt aat cca ctc act ttc ggt gct ggg acc aag ctg gag ctg aaa cgg Ser Asn Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys Arg	780
225 230 235	
gcg gcc gca gaacaaaaac tcattctcaga agaggatctg aatggatcca	829
Ala Ala Ala 240	
aagacgaact ctag	843

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